

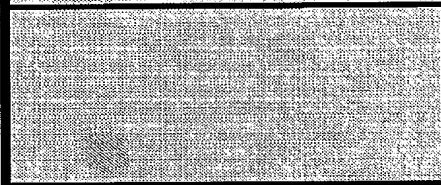



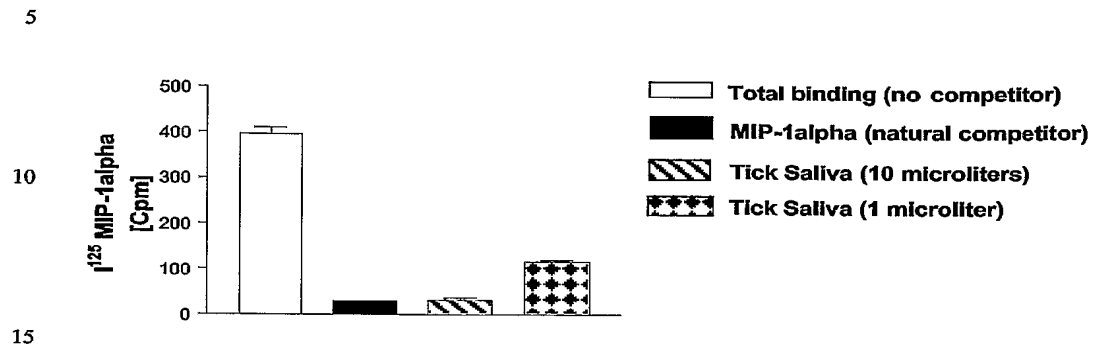
Figure 1

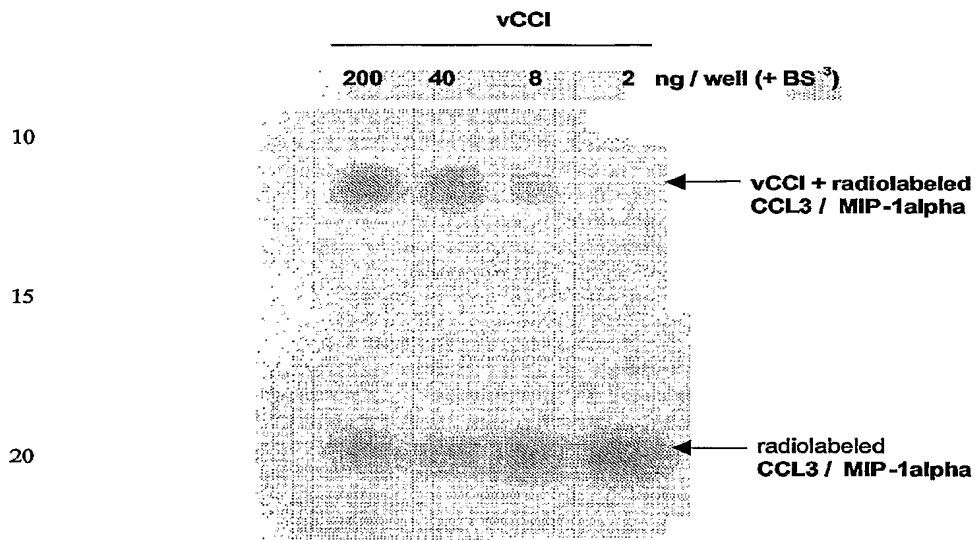
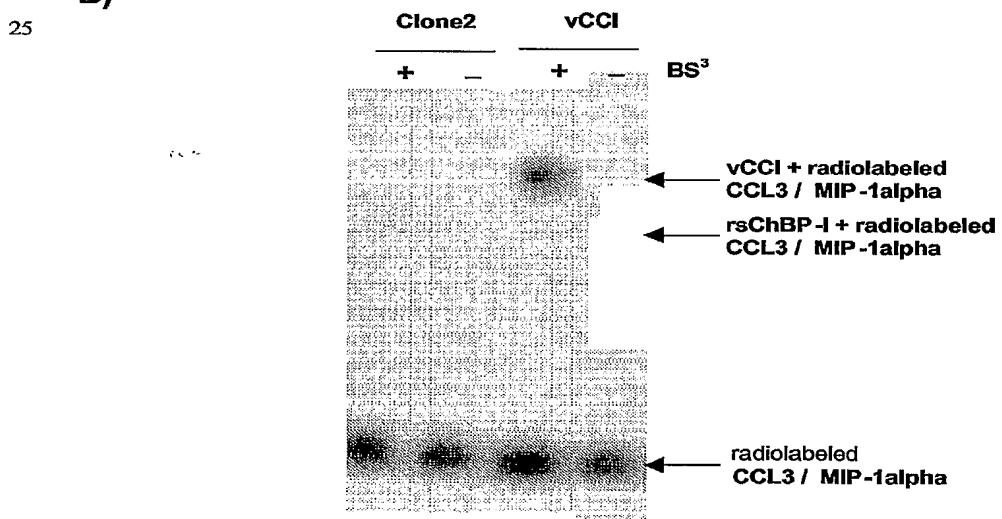
5

Spotted Filter

10

2 100	0.2 10	0.1 1	microliters rhSE µg vCCI	Radiolabeled Chemokine
				CCL2 / MCP-1
				CCL3 / MIP-1alpha
				CCL5 / RANTES
				CXCL8 / Interleukin-8

**Figure 2**

**Figure 3**5 **A)**25 **B)**

**Figure 4**

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5      1  GGCCATTACGGCCGGGGTCCTTGCGCATTTCGTGTAGAGCAGCAGCTCAAGTCTTCGAAG

        61  ATG  CGC  ACT  TTC  GGG  GCT  TCT  CTT  TTC  GTT
          1  Met  Arg  Thr  Phe  Gly  Ala  Ser  Leu  Phe  Val

10      91  CTC  CTC  GCG  ATT  AGT  GTC  GCT  TAC  TGT  GAC
        11  Leu  Leu  Ala  Ile  Ser  Val  Ala  Tyr  Cys  Asp

        121  GTC  CAA  GAG  CGC  GGC  CAT  ACT  TAC  GTG  ACC
          21  Val  Gln  Glu  Arg  Gly  His  Thr  Tyr  Val  Thr

15      151  AAA  AAT  GTG  ACG  GTC  GAA  AAC  GGT  GCC  TGC
          31  Lys  Asn  Val  Thr  Val  Glu  Asn  Gly  Ala  Cys

        181  GTG  TTT  GAA  CGC  AAC  GTC  ATT  CCG  GAT  GGT
        41  Val  Phe  Glu  Arg  Asn  Val  Ile  Pro  Asp  Gly

        210  GAA  ACC  AAA  GCA  CTG  AAC  AGC  CCA  TGC  GTC
          51  Glu  Thr  Lys  Ala  Leu  Asn  Ser  Pro  Cys  Val

20      241  ATT  TCC  ACA  TGC  TAT  GCA  GCT  GAC  CGT  AAA
          61  Ile  Ser  Thr  Cys  Tyr  Ala  Ala  Asp  Arg  Lys

        271  GTG  AAC  TCG  ACT  CTC  TGC  CCG  AAC  TTC  GGA
          71  Val  Asn  Ser  Thr  Leu  Cys  Pro  Asn  Phe  Gly

30      301  GTT  GCG  GAG  GGC  TGC  CAT  GTG  GAG  TGG  ACC
          81  Val  Ala  Glu  Gly  Cys  His  Val  Glu  Trp  Thr

        331  CCC  GAT  GGT  GAA  TAC  CCG  AAC  TGC  TGC  CCG
        91  Pro  Asp  Gly  Glu  Tyr  Pro  Asn  Cys  Cys  Pro

35      361  AAG  CAT  GTG  TGC  CCT  ACG  GCC  CCT  GTT  ACT
        101  Lys  His  Val  Cys  Pro  Thr  Ala  Pro  Val  Thr

40      391  TCT  TAA  TCGCATCACATCTGCGAAAATGAA
        111  Ser  STOP

        421  ACGTCGAGACATTCTTCTTTATGCCTTAAGAAATTAAACTGCAACGTCCGCAAAAATACA

45      481  TCCCCGCTTCAAATACGAACAAAATGCAGGATCAAATGCTATTAGGTTTCATGCTGA  GTG

        541  CAAGCTAAATAAACTGAATCAGCGTTTAAAAAAAAAAAAAAAA

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**Figure 5**

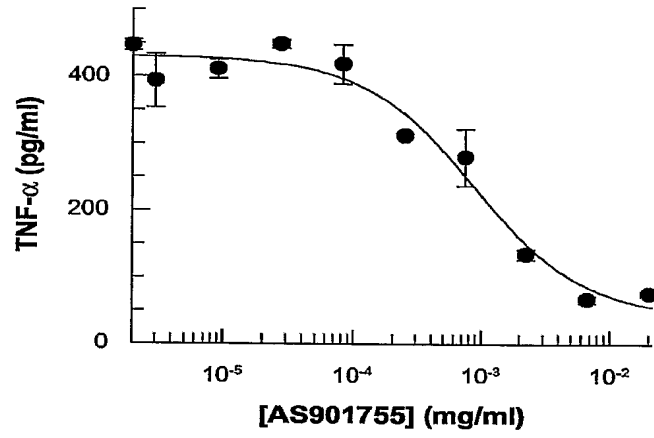
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5  avChBP-I  60  MRALVALACVVSVAVVIGDIQEHGHSYLKRNVTIENGACIYERNITLPDGETKALHDPCV 239
    MR  A  V++++V  D+QE GH+Y+ +NVT+ENGAC++ERN +PDGETKAL+ PCV
rsChBP-I  61  MRTEGASLFVLLAISVAY CDVQERGHYVTKNVTVENGACVFERNVIPDGETKALNSPCV 240
    MR+  + L  H Y  V+++NG C +  + DG++  + PC
10 isChBP-I  1  MRSIVLWALIALGGVPLLGMGAANQSHPYG ---VSFNNGTCTYRNITLRDGDSEFFQYPCE 171

avChBP-I  240  IATCYAERREVNATLCPNFGVDPGCRVQWTPDGIYPECCPKQV CDGTN 383
    I+TCYA R VN+TLCENFGV GC V WTEPDG YP CCEK VC
15 rsChBP-I  241  ISTCYAADRKVNSTLCPNFGVAEGCHEVWTPDGEYFNCCPKHVCPTAPVTS 393
    C R + C C +P CC
isChBP-I  172  YWNCNVTARTLTIEGCG -VPYGSCLYVHNYNFYWPLCCRMSRLC 303

20

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**Figure 6**

**Figure 7**